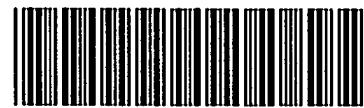


RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/889,330
Source: JFw16
Date Processed by STIC: 1/14/05

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IFW16

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/889,330

DATE: 01/14/2005
TIME: 13:00:17

Input Set : A:\249146us.app
Output Set: N:\CRF4\01142005\I889330.raw

3 <110> APPLICANT: KOLTERMAN, ORVILLE G.
 4 YOUNG, ANDREW A.
 6 <120> TITLE OF INVENTION: NOVEL EXENDIN AGONIST FORMULATIONS AND METHODS OF
 ADMINISTRATION
 7 THEREOF
 9 <130> FILE REFERENCE: 249/146US
 11 <140> CURRENT APPLICATION NUMBER: 09/889,330
 12 <141> CURRENT FILING DATE: 2001-12-27
 14 <150> PRIOR APPLICATION NUMBER: PCT/US00/00902
 15 <151> PRIOR FILING DATE: 2000-01-14
 17 <150> PRIOR APPLICATION NUMBER: US 60/116,380
 18 <151> PRIOR FILING DATE: 1999-01-14
 20 <150> PRIOR APPLICATION NUMBER: US 60/175,365
 21 <151> PRIOR FILING DATE: 2000-01-10
 23 <160> NUMBER OF SEQ ID NOS: 189
 25 <170> SOFTWARE: PatentIn Ver. 3.2
 27 <210> SEQ ID NO: 1
 28 <211> LENGTH: 39
 29 <212> TYPE: PRT
 30 <213> ORGANISM: Heloderma horridum
 32 <220> FEATURE:
 33 <221> NAME/KEY: MOD_RES
 34 <222> LOCATION: (39)
 35 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2
 37 <400> SEQUENCE: 1
 38 His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 39 1 5 10 15
 41 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 42 20 25 30
 44 Ser Gly Ala Pro Pro Pro Ser
 45 35
 48 <210> SEQ ID NO: 2
 49 <211> LENGTH: 39
 50 <212> TYPE: PRT
 51 <213> ORGANISM: Heloderma suspectum
 53 <220> FEATURE:
 54 <221> NAME/KEY: MOD_RES
 55 <222> LOCATION: (39)
 56 <223> OTHER INFORMATION: AMIDATION, Position 39 is Ser-NH2
 58 <400> SEQUENCE: 2
 59 His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
 60 1 5 10 15
 62 Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
 63 20 25 30

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RAW SEQUENCE LISTING
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Input Set : A:\249146us.app
Output Set: N:\CRF4\01142005\I889330.raw

65 Ser Gly Ala Pro Pro Pro Ser
66 35
69 <210> SEQ ID NO: 3
70 <211> LENGTH: 39
71 <212> TYPE: PRT
72 <213> ORGANISM: Artificial Sequence
74 <220> FEATURE:
75 <223> OTHER INFORMATION: synthetic construct
77 <220> FEATURE:
78 <221> NAME/KEY: VARIANT
79 <222> LOCATION: (1) /
80 <223> OTHER INFORMATION: His, Arg or Thr
82 <220> FEATURE:
83 <221> NAME/KEY: VARIANT /
84 <222> LOCATION: (2) /
85 <223> OTHER INFORMATION: Ser, Gly, Ala or Thr
87 <220> FEATURE:
88 <221> NAME/KEY: VARIANT /
89 <222> LOCATION: (3) /
90 <223> OTHER INFORMATION: Asp or Glu
92 <220> FEATURE:
93 <221> NAME/KEY: VARIANT /
94 <222> LOCATION: (6) /
95 <223> OTHER INFORMATION: Phe, Tyr or naphthalanine
97 <220> FEATURE:
98 <221> NAME/KEY: VARIANT /
99 <222> LOCATION: (7) /
100 <223> OTHER INFORMATION: Thr or Ser
102 <220> FEATURE:
103 <221> NAME/KEY: VARIANT /
104 <222> LOCATION: (8) /
105 <223> OTHER INFORMATION: Ser or Thr
107 <220> FEATURE:
108 <221> NAME/KEY: VARIANT /
109 <222> LOCATION: (9) /
110 <223> OTHER INFORMATION: Asp or Glu
112 <220> FEATURE:
113 <221> NAME/KEY: VARIANT /
114 <222> LOCATION: (10) /
115 <223> OTHER INFORMATION: Leu, Ile, Val, pentylglycine or Met
117 <220> FEATURE:
118 <221> NAME/KEY: VARIANT /
119 <222> LOCATION: (14) /
120 <223> OTHER INFORMATION: Leu, Ile, pentylglycine, Val or Met
122 <220> FEATURE:
123 <221> NAME/KEY: VARIANT /
124 <222> LOCATION: (22) /
125 <223> OTHER INFORMATION: Phe, Tyr or naphthalanine
127 <220> FEATURE:

RAW SEQUENCE LISTING

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Input Set : A:\249146us.app

Output Set: N:\CRF4\01142005\I889330.raw

128 <221> NAME/KEY: VARIANT
 129 <222> LOCATION: (23)
 130 <223> OTHER INFORMATION: Ile, Val, Leu, pentylglycine, tert-butylglycine or Met
 132 <220> FEATURE:
 133 <221> NAME/KEY: VARIANT
 134 <222> LOCATION: (24)
 135 <223> OTHER INFORMATION: Glu or Asp
 137 <220> FEATURE:
 138 <221> NAME/KEY: VARIANT
 139 <222> LOCATION: (25)
 140 <223> OTHER INFORMATION: Trp, Phe, Tyr or naphthylalanine
 142 <220> FEATURE:
 143 <221> NAME/KEY: VARIANT
 144 <222> LOCATION: (31)
 145 <223> OTHER INFORMATION: independently Pro, homoproline, 3-hydroxyproline,
 146 4-hydroxyproline, thioproline, N-alkylglycine,
 147 N-alkylpentylglycine or N-alkylalanine
 149 <220> FEATURE:
 150 <221> NAME/KEY: VARIANT
 151 <222> LOCATION: (36)..(38)
 152 <223> OTHER INFORMATION: independently Pro, homoproline, 3-hydroxyproline,
 153 4-hydroxyproline, thioproline, N-alkylglycine,
 154 N-alkylpentylglycine or N-alkylalanine
 156 <220> FEATURE:
 157 <221> NAME/KEY: VARIANT
 158 <222> LOCATION: (39)
 159 <223> OTHER INFORMATION: Ser, Thr or Tyr
 161 <220> FEATURE:
 162 <223> OTHER INFORMATION: c-term is -OH or NH₂, with the proviso that the compound
 163 does not have the formula of either SEQ ID NOS 1 or 2
 165 <400> SEQUENCE: 3
 W--> 166 Xaa Xaa Xaa Gly Thr Xaa Xaa Xaa Xaa Ser Lys Gln Xaa Glu Glu
 167 1 5 10 15
 169 Glu Ala Val Arg Leu Xaa Xaa Xaa Xaa Leu Lys Asn Gly Gly Xaa Ser
 170 20 25 30
 172 Ser Gly Ala Xaa Xaa Xaa Xaa
 173 35
 175 <210> SEQ ID NO: 4
 176 <211> LENGTH: 38
 177 <212> TYPE: PRT
 178 <213> ORGANISM: Artificial Sequence
 180 <220> FEATURE:
 181 <223> OTHER INFORMATION: synthetic construct
 183 <220> FEATURE:
 184 <221> NAME/KEY: VARIANT
 185 <222> LOCATION: (1)
 186 <223> OTHER INFORMATION: His, Arg, or Tyr
 188 <220> FEATURE:
 189 <221> NAME/KEY: VARIANT

RAW SEQUENCE LISTING
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DATE: 01/14/2005
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Input Set : A:\249146us.app
Output Set: N:\CRF4\01142005\I889330.raw

190 <222> LOCATION: (2)
191 <223> OTHER INFORMATION: Ser, Gly, Ala or Thr
193 <220> FEATURE:
194 <221> NAME/KEY: VARIANT
195 <222> LOCATION: (3)
196 <223> OTHER INFORMATION: Asp or Glu
198 <220> FEATURE:
199 <221> NAME/KEY: VARIANT
200 <222> LOCATION: (5)
201 <223> OTHER INFORMATION: Ala or Thr
203 <220> FEATURE:
204 <221> NAME/KEY: VARIANT
205 <222> LOCATION: (6)
206 <223> OTHER INFORMATION: Ala, Phe, Tyr or naphthalalanine
208 <220> FEATURE:
209 <221> NAME/KEY: VARIANT
210 <222> LOCATION: (7)
211 <223> OTHER INFORMATION: Thr or Ser
213 <220> FEATURE:
214 <221> NAME/KEY: VARIANT
215 <222> LOCATION: (8)
216 <223> OTHER INFORMATION: Ala, Ser or Thr
218 <220> FEATURE:
219 <221> NAME/KEY: VARIANT
220 <222> LOCATION: (9)
221 <223> OTHER INFORMATION: Asp or Glu
223 <220> FEATURE:
224 <221> NAME/KEY: VARIANT
225 <222> LOCATION: (10)
226 <223> OTHER INFORMATION: Ala, Leu, Ile, Val, pentylglycine or Met
228 <220> FEATURE:
229 <221> NAME/KEY: VARIANT
230 <222> LOCATION: (11)
231 <223> OTHER INFORMATION: Ala or Ser
233 <220> FEATURE:
234 <221> NAME/KEY: VARIANT
235 <222> LOCATION: (12)
236 <223> OTHER INFORMATION: Ala or Lys
238 <220> FEATURE:
239 <221> NAME/KEY: VARIANT
240 <222> LOCATION: (13)
241 <223> OTHER INFORMATION: Ala or Gln
243 <220> FEATURE:
244 <221> NAME/KEY: VARIANT
245 <222> LOCATION: (14)
246 <223> OTHER INFORMATION: Ala, Leu, Ile, pentylglycine, Val or Met
248 <220> FEATURE:
249 <221> NAME/KEY: VARIANT
250 <222> LOCATION: (15)..(17)

RAW SEQUENCE LISTING
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DATE: 01/14/2005
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Input Set : A:\249146us.app
Output Set: N:\CRF4\01142005\I889330.raw

251 <223> OTHER INFORMATION: Ala or Glu
253 <220> FEATURE:
254 <221> NAME/KEY: VARIANT
255 <222> LOCATION: (19)
256 <223> OTHER INFORMATION: Ala or Val
258 <220> FEATURE:
259 <221> NAME/KEY: VARIANT
260 <222> LOCATION: (20)
261 <223> OTHER INFORMATION: Ala or Arg
263 <220> FEATURE:
264 <221> NAME/KEY: VARIANT
265 <222> LOCATION: (21)
266 <223> OTHER INFORMATION: Ala or Leu
268 <220> FEATURE:
269 <221> NAME/KEY: VARIANT
270 <222> LOCATION: (22)
271 <223> OTHER INFORMATION: Ala, Phe, Tyr or naphthylalanine
273 <220> FEATURE:
274 <221> NAME/KEY: VARIANT
275 <222> LOCATION: (23)
276 <223> OTHER INFORMATION: Ile, Val, Leu, pentylglycine, tert-butylglycine or Met
278 <220> FEATURE:
279 <221> NAME/KEY: VARIANT
280 <222> LOCATION: (24)
281 <223> OTHER INFORMATION: Ala, Glu or Asp
283 <220> FEATURE:
284 <221> NAME/KEY: VARIANT
285 <222> LOCATION: (25)
286 <223> OTHER INFORMATION: Ala, Trp, Phe, Tyr or naphthylalanine
288 <220> FEATURE:
289 <221> NAME/KEY: VARIANT
290 <222> LOCATION: (26)
291 <223> OTHER INFORMATION: Ala or Leu
293 <220> FEATURE:
294 <221> NAME/KEY: VARIANT
295 <222> LOCATION: (27)
296 <223> OTHER INFORMATION: Ala or Lys
298 <220> FEATURE:
299 <221> NAME/KEY: VARIANT
300 <222> LOCATION: (28)
301 <223> OTHER INFORMATION: Ala or Asn
303 <220> FEATURE:
304 <221> NAME/KEY: MOD_RES
305 <222> LOCATION: (31)
306 <223> OTHER INFORMATION: Pro, homoproline, 3Hyp, 4Hyp, thioproline, N-alkylglycine,
307 N-alkylpentylglycine or N-alkylalanine
309 <220> FEATURE:
310 <221> NAME/KEY: MOD_RES
311 <222> LOCATION: (36)..(38)

RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : A:\249146us.app
Output Set: N:\CRF4\01142005\I889330.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 1,2,3,5,7,8,9,10,14,22,23,24,25,31,36,37,38,39
Seq#:4; Xaa Pos. 1,2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23,24
Seq#:4; Xaa Pos. 25,26,27,28,31,36,37,38
Seq#:5; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,19,20,21,22,23
Seq#:5; Xaa Pos. 24,25,26,27,28,31,36,37,38,39
Seq#:15; Xaa Pos. 6
Seq#:20; Xaa Pos. 10
Seq#:21; Xaa Pos. 10
Seq#:22; Xaa Pos. 14
Seq#:23; Xaa Pos. 14
Seq#:24; Xaa Pos. 22
Seq#:27; Xaa Pos. 23
Seq#:28; Xaa Pos. 23
Seq#:31; Xaa Pos. 31,36,37,38
Seq#:32; Xaa Pos. 36,37,38
Seq#:33; Xaa Pos. 31,36,37,38
Seq#:34; Xaa Pos. 36,37,38
Seq#:35; Xaa Pos. 31,36,37,38
Seq#:36; Xaa Pos. 31,36,37,38
Seq#:37; Xaa Pos. 31,36,37,38
Seq#:38; Xaa Pos. 36,37,38
Seq#:39; Xaa Pos. 31,36,37,38
Seq#:81; Xaa Pos. 31,36,37,38
Seq#:82; Xaa Pos. 36,37,38
Seq#:83; Xaa Pos. 31
Seq#:84; Xaa Pos. 31,36,37
Seq#:85; Xaa Pos. 31,36,37
Seq#:86; Xaa Pos. 31,36
Seq#:89; Xaa Pos. 6
Seq#:93; Xaa Pos. 10
Seq#:94; Xaa Pos. 22
Seq#:95; Xaa Pos. 23
Seq#:99; Xaa Pos. 31,36,37
Seq#:115; Xaa Pos. 6
Seq#:116; Xaa Pos. 6
Seq#:127; Xaa Pos. 10
Seq#:128; Xaa Pos. 10
Seq#:137; Xaa Pos. 14
Seq#:138; Xaa Pos. 14
Seq#:151; Xaa Pos. 22
Seq#:152; Xaa Pos. 22
Seq#:155; Xaa Pos. 23
Seq#:156; Xaa Pos. 23
Seq#:181; Xaa Pos. 31,36,37,38

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Seq#:182; Xaa Pos. 36,37,38
Seq#:183; Xaa Pos. 31,36,37
Seq#:184; Xaa Pos. 31,36

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:5; Line(s) 484

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Input Set : A:\249146us.app
Output Set: N:\CRF4\01142005\I889330.raw

L:166 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
M:341 Repeated in SeqNo=3
L:320 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
M:341 Repeated in SeqNo=4
L:481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
M:341 Repeated in SeqNo=5
L:711 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:836 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:894 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:16
L:1032 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:16
L:1061 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:16
L:1143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:16
M:341 Repeated in SeqNo=31
L:1175 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:32
L:1206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:16
M:341 Repeated in SeqNo=33
L:1238 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:32
L:1269 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:16
M:341 Repeated in SeqNo=35
L:1303 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:16
M:341 Repeated in SeqNo=36
L:1337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:16
M:341 Repeated in SeqNo=37
L:1369 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:32
L:1400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:16
M:341 Repeated in SeqNo=39
L:2331 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:16
M:341 Repeated in SeqNo=81
L:2363 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 after pos.:32
L:2389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:16
L:2423 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 after pos.:16
M:341 Repeated in SeqNo=84
L:2457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:16
M:341 Repeated in SeqNo=85
L:2491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:16
M:341 Repeated in SeqNo=86
L:2562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:89 after pos.:0
L:2651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93 after pos.:0
L:2680 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:16
L:2706 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95 after pos.:16
L:2803 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:99 after pos.:16
M:341 Repeated in SeqNo=99
L:3144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:115 after pos.:0
L:3170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:116 after pos.:0
L:3406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:127 after pos.:0

VERIFICATION SUMMARY

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L:3432 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:128 after pos.:0

L:3626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:137 after pos.:0